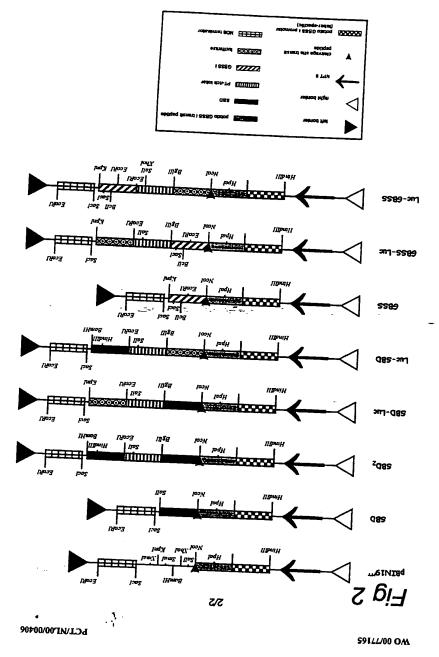
FIG 7 Alignment of starch-binding domains

maltose-binding site 1 📕 maltose-binding site 2 📕 disulfide bridge	G-STVTEEGGSNHTETIAPSSGTATINVNWQ-P SDDSVEESDPNREYTVPQA GTSTATVTDTWR G-NTITEGGSNHTYTVPSSSTGTVIVNWQQ SQGNVTEESGSNHVYTTPTNTTGKIIVDWQN	AKAIIGPMYNQVVYQYENWYYDVSVEAGKTIEFKFLKKQ SDGIALSADKITSSDELWYVTVTLEAGESFEYKFIRIE SKAIIGPMFNQVVYQYETWYYDVSVEAGTTIQFKFIKKD SKAIIGPMFNQVVYSYETWYIDVSVEEGKTIEFKFIKKD	SGDQVSVREVVNNATTALGQNVYLTGSVSELGNEDP TTPTAVAVTEDLT-ATTT GENIYLVGSISQLGD ET TGNQICVREVVNNASTVYGENVYLTGNVAELGN DT TNDQVSVREVVNNATTNLGQNIYIVGNVYELGN DT
	CGT - B. circulans GA - A. niger CGT - T. thermosulfurigenes CGT - B. stearothermophilus	CGT - B. circulans GA - A. niger CGT - T. thermosulfurigenes CGT - B. stearothermophilus	CGT - B. circulans GA - A. niger CGT - T. thermosulfurlgenes CGT - B. stearothermophilus
		1/2	

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